



1627 ENTERED

1600

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/726,219

DATE: 03/13/2002 P.5
TIME: 14:58:16

Input Set : A:\50020075_1.RTF
Output Set: N:\CRF3\03132002\I726219.raw

1600
MAR 22 2002
FBI/CA/EN 1000/2000

3 <110> APPLICANT: Cambridge Antibody Technology
4 Cambridge Antibody Technology Limited
5 Medical Research Council
6 McCafferty, John
7 Pope, Anthony
8 Johnson, Kevin
9 Hoogenboom, Hendricus
10 Griffiths, Andrew
11 Jackson, Ronald
12 Holliger, Kasper
13 Marks, James
14 Clackson, Timothy
15 Chiswell, David
16 Winter, Gregory
17 Bonert, Timothy
19 <120> TITLE OF INVENTION: Methods for Producing Members of Specific Binding Pairs
21 <130> FILE REFERENCE: 13839-00013
23 <140> CURRENT APPLICATION NUMBER: US 09/726,219
24 <141> CURRENT FILING DATE: 2000-11-28
26 <150> PRIOR APPLICATION NUMBER: GB 9015198.6
27 <151> PRIOR FILING DATE: 1990-07-10
29 <150> PRIOR APPLICATION NUMBER: GB 9022845.3
30 <151> PRIOR FILING DATE: 1990-10-19
W--> 32 <150> PRIOR APPLICATION NO: GB 9022845.3
33 <151> PRIOR FILING DATE: 1990-10-19
35 <150> PRIOR APPLICATION NUMBER: GB 9024503.6
36 <151> PRIOR FILING DATE: 1990-11-12
38 <150> PRIOR APPLICATION NUMBER: GB 9104744.9
39 <151> PRIOR FILING DATE: 1991-03-06
41 <150> PRIOR APPLICATION NUMBER: GB 9110549.4
42 <151> PRIOR FILING DATE: 1991-05-15
44 <150> PRIOR APPLICATION NUMBER: PCT/GB91/01134
45 <151> PRIOR FILING DATE: 1991-07-10
47 <150> PRIOR APPLICATION NUMBER: US 07/971,857
48 <151> PRIOR FILING DATE: 1993-01-08
50 <150> PRIOR APPLICATION NUMBER: US 08/484,893
51 <151> PRIOR FILING DATE: 1995-06-07
53 <160> NUMBER OF SEQ ID NOS: 272
55 <170> SOFTWARE: PatentIn version 3.1
57 <210> SEQ ID NO: 1
58 <211> LENGTH: 5
59 <212> TYPE: PRT
60 <213> ORGANISM: Bacteriophage fd

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65 1 5
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69 <211> LENGTH: 5
70 <212> TYPE: PRT
71 <213> ORGANISM: Bacteriophage fd
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76 1 5
79 <210> SEQ ID NO: 3
80 <211> LENGTH: 5
81 <212> TYPE: PRT
82 <213> ORGANISM: Bacteriophage fd
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86 Leu Glu Ile Lys Arg
87 1 5
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91 <211> LENGTH: 75
92 <212> TYPE: DNA
93 <213> ORGANISM: Artificial Sequence
95 <220> FEATURE:
96 <223> OTHER INFORMATION: oligonucleotide for mutagenesis
98 <400> SEQUENCE: 4
99 actttcaaca gtttctgcgg ccgcccgttt gatctcgagc tcctgcagtt ggacctgtgc 60
101 actgtgagaa tagaa 75
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105 <211> LENGTH: 22
106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: PCR primer
112 <400> SEQUENCE: 5
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116 <210> SEQ ID NO: 6
117 <211> LENGTH: 34
118 <212> TYPE: DNA
119 <213> ORGANISM: Artificial Sequence
121 <220> FEATURE:
122 <223> OTHER INFORMATION: PCR primer
124 <400> SEQUENCE: 6
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129 <211> LENGTH: 27
130 <212> TYPE: DNA
131 <213> ORGANISM: Artificial Sequence
133 <220> FEATURE:
134 <223> OTHER INFORMATION: PCR primer
136 <400> SEQUENCE: 7

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141 <211> LENGTH: 24

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143 <213> ORGANISM: Artificial Sequence

145 <220> FEATURE:

146 <223> OTHER INFORMATION: oligonucleotide probe distinguishing between pAb D1.3 and

pAB NQ1

147 1

149 <400> SEQUENCE: 8

150 gtagtcaagc ctataatctc tctc 24

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154 <211> LENGTH: 51

155 <212> TYPE: DNA

156 <213> ORGANISM: Artificial Sequence

158 <220> FEATURE:

159 <223> OTHER INFORMATION: PCR primer

161 <400> SEQUENCE: 9

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167 <212> TYPE: DNA

168 <213> ORGANISM: Artificial Sequence

170 <220> FEATURE:

171 <223> OTHER INFORMATION: PCR primer

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178 <211> LENGTH: 33

179 <212> TYPE: DNA

180 <213> ORGANISM: Artificial Sequence

182 <220> FEATURE:

183 <223> OTHER INFORMATION: PCR primer

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191 <212> TYPE: DNA

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194 <220> FEATURE:

195 <223> OTHER INFORMATION: PCR primer

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202 <211> LENGTH: 24

203 <212> TYPE: DNA

204 <213> ORGANISM: Artificial Sequence

206 <220> FEATURE:

207 <223> OTHER INFORMATION: oligonucleotide for mutagenesis - removal of a BamH1 site

209 <400> SEQUENCE: 13

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214 <211> LENGTH: 26

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218 <220> FEATURE:

219 <223> OTHER INFORMATION: oligonucleotide for mutagenesis - introduction of a BamH1
site

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226 <211> LENGTH: 15

227 <212> TYPE: PRT

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230 <220> FEATURE:

231 <223> OTHER INFORMATION: linker between VH and VLK

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236 1 5 10 15

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242 <213> ORGANISM: Artificial Sequence

244 <220> FEATURE:

245 <223> OTHER INFORMATION: primer for reverse transcription

247 <400> SEQUENCE: 16

248 ctggacaggg atccagagtt cca 23

251 <210> SEQ ID NO: 17

252 <211> LENGTH: 23

253 <212> TYPE: DNA

254 <213> ORGANISM: Artificial Sequence

256 <220> FEATURE:

257 <223> OTHER INFORMATION: primer for reverse transcription

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266 <213> ORGANISM: Artificial Sequence

268 <220> FEATURE:

269 <223> OTHER INFORMATION: PCR primer

271 <400> SEQUENCE: 18

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275 <210> SEQ ID NO: 19

276 <211> LENGTH: 22

277 <212> TYPE: DNA

278 <213> ORGANISM: Artificial Sequence

280 <220> FEATURE:

281 <223> OTHER INFORMATION: PCR primer

283 <400> SEQUENCE: 19

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Input Set : A:\50020075_1.RTF

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295 <400> SEQUENCE: 20
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300 <211> LENGTH: 24
301 <212> TYPE: DNA
302 <213> ORGANISM: Artificial Sequence
304 <220> FEATURE:
305 <223> OTHER INFORMATION: PCR primer
307 <400> SEQUENCE: 21
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311 <210> SEQ ID NO: 22
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313 <212> TYPE: DNA
314 <213> ORGANISM: Artificial Sequence
316 <220> FEATURE:
317 <223> OTHER INFORMATION: PCR primer
319 <400> SEQUENCE: 22
320 ccgttttatt tccaactttg tccc                                24
323 <210> SEQ ID NO: 23
324 <211> LENGTH: 24
325 <212> TYPE: DNA
326 <213> ORGANISM: Artificial Sequence
328 <220> FEATURE:
329 <223> OTHER INFORMATION: PCR primer
331 <400> SEQUENCE: 23
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335 <210> SEQ ID NO: 24
336 <211> LENGTH: 24
337 <212> TYPE: DNA
338 <213> ORGANISM: Artificial Sequence
340 <220> FEATURE:
341 <223> OTHER INFORMATION: PCR primer
343 <400> SEQUENCE: 24
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347 <210> SEQ ID NO: 25
348 <211> LENGTH: 24
349 <212> TYPE: DNA
350 <213> ORGANISM: Artificial Sequence
352 <220> FEATURE:
353 <223> OTHER INFORMATION: PCR primer
355 <400> SEQUENCE: 25
356 tggagactcg gtgagctcaa tgtc                                24

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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

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Input Set : A:\50020075_1.RTF

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L:32 M:288 W: Application Number is Repeated, <150> PRIOR APPLICATION NUMBER
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:995 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75
L:1014 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76
L:1033 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77